**FIG. 1**

CG ACGGCCCGGC TGGTAAATTC CCCTTTCTCC  
 -690 AAAATGTAAA ATAAATCTGC TTCCATCTTC TAAATACTA TGGGACTAAA  
 -640 CATCCTTTTG TTATGCTAAG GAAAAGCCAG TATTCGCGTT GATTTAGAAG  
 -590 AGGGATGTTC TGGTTATAGA ACGATGCTGT GTCTCAGAAA CACTTAAATA  
 -540 CTATTAAGCT AGAAATAGAA GGGAAAATAA TGCTTCCCCG CATCTCCCCT  
 -490 CAAGTGTAGT CCTCTTTTTT TAGCCTGATT TCCGACGAAA TGTCTGAATG  
 -440 CCTACAGTTA TTTGGCCATC CTGAAAAGTG CAACTTATCC TGACGTCTCG

## CRE

-390 AGGGACGGAA AAGTTACCGA AGTCCAAGGA ATGAGTCACT TTGCTCAAAT  
 -340 TTGATGAGTA ATATCAGGTG TCATGAAACC CAGTTTCGAA GGAGAGGGGA  
 -290 GGGGGCGTCA GATCTGCAGA CGGAAGCAGG CCGCTCCGGA TTGGATGGCG  
 -240 AGACCTCGAT TTTCTTAAAA TTGCGTCATT TAGAACCCAA TTGGGTCCAG

## CRE-like

-190 ATGTTATGGG CATCGACGAG TTACCGTCTC GGAAACTCTC AATCAGCAA  
 -140 GCGAAAGGAG AGGAGGCGGC TAATTAAATA TTGAGCAGAA AGTCGCGTGG  
 -90 GGAGAATGTC ACGTGGGTCT GGAGGCTCAA GGAGGCTGGG ATAAATACCG  
 -40 CAAGGCACTG AGCAGGCGAA AGAGCGCGCT CGGACCTCCT  
 +1 TTCCCGGCGG CAGCTACCGA GAGTGCGGAG CGACCAGCGT GCGCTCGGAG

## Exon 1

+51 AACCAGAGAA CTCAGCACCC CGCGGGACTG TCCGTCGCAG TAAGTGCCCG

## Intron 1

+101 CGCGGTGCTG GCCGCGGCTG CCCGGGTCAT CCCACCCCGC ATCTGTCCGA  
 +151 GGTGGCCGCG CTGGGGGCGC CGCTGCGGCG AGGGACAGTG GGGAGACTGG  
 +201 CTTCCCAAAC CCCAACCACC CTCTTTGTCT TCCACCTGCA GAGTTTCCTG  
 +251 GTTTGAAGGT GTGGGTTGGT GGGTTAGGGG GCTGGGGGAG CTGGGATTCA  
 +301 GGGAGAAGAG GGTGGAGAA TCTTTGGGAC GCGATTCTCT CGCCTAACCG  
 +351 GTACAGGTGA GACTTCAGTC CTTATGTTTT TGATCTTGGT TCATCCGTTG  
 +401 TGGGGCAGAA AATTCTGTTG CTTTAACTCT TGGATAACCA CCCCTAATAG  
 +451 ATACATTATT TCTCTCTTTG GTGTCTTCTC CTCCTACCCC TTCCAGAAA

## Exon 2

+501 TCCGAC

FIG. 2

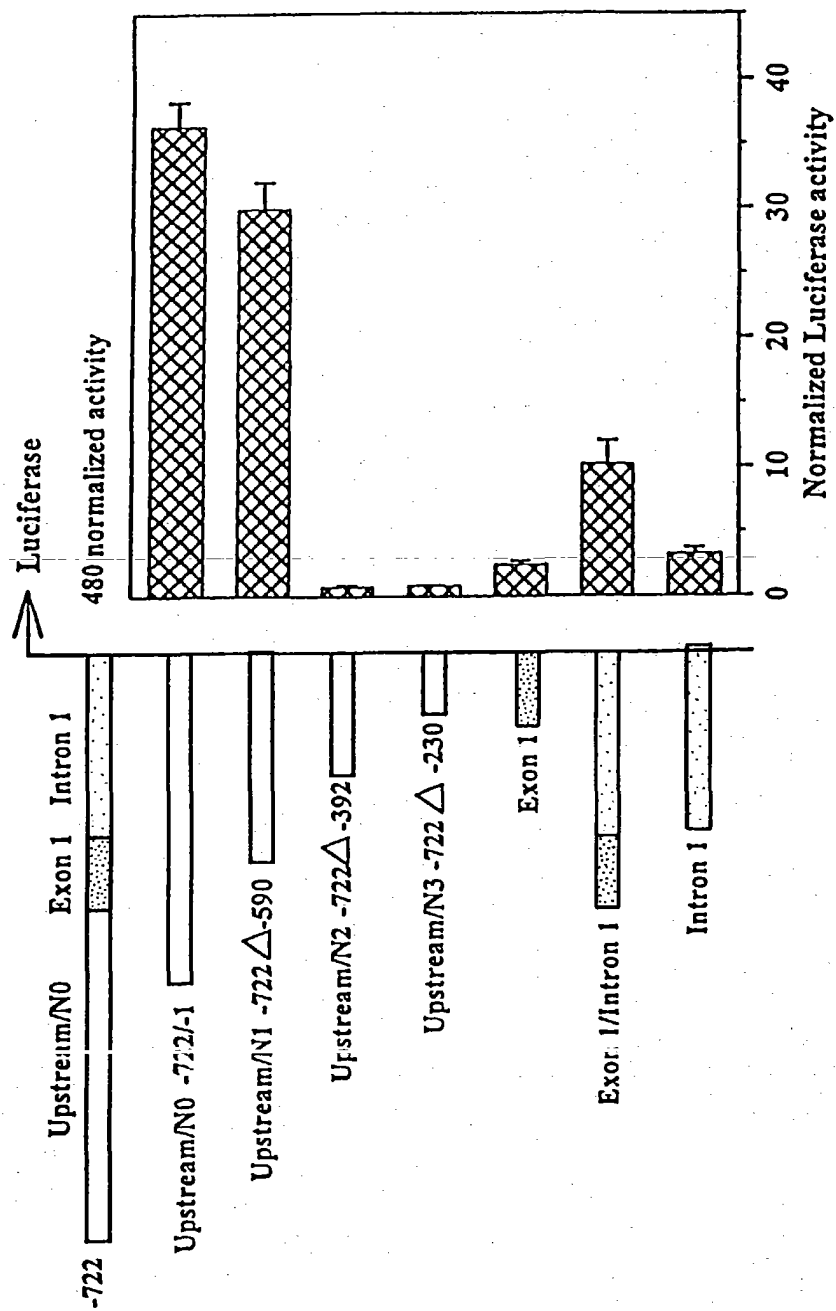


FIG. 3

FIG. 4A

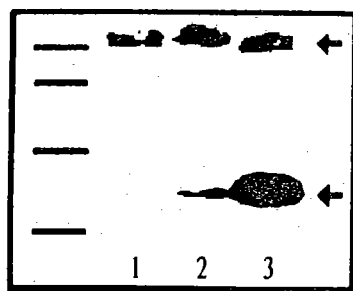
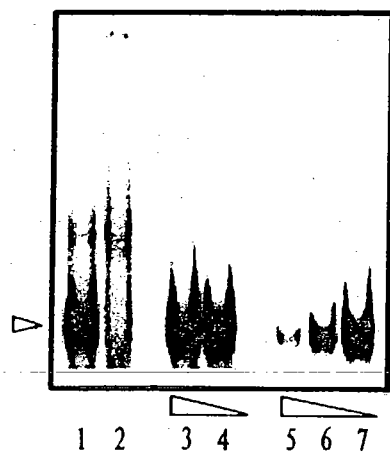


FIG. 4B

FIG. 4C

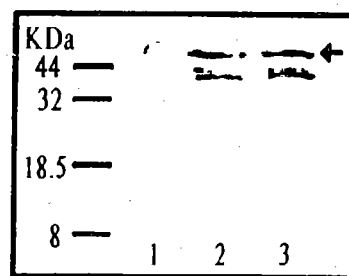
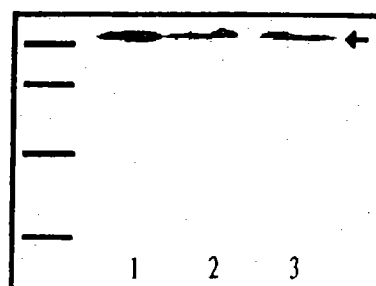


FIG. 4D

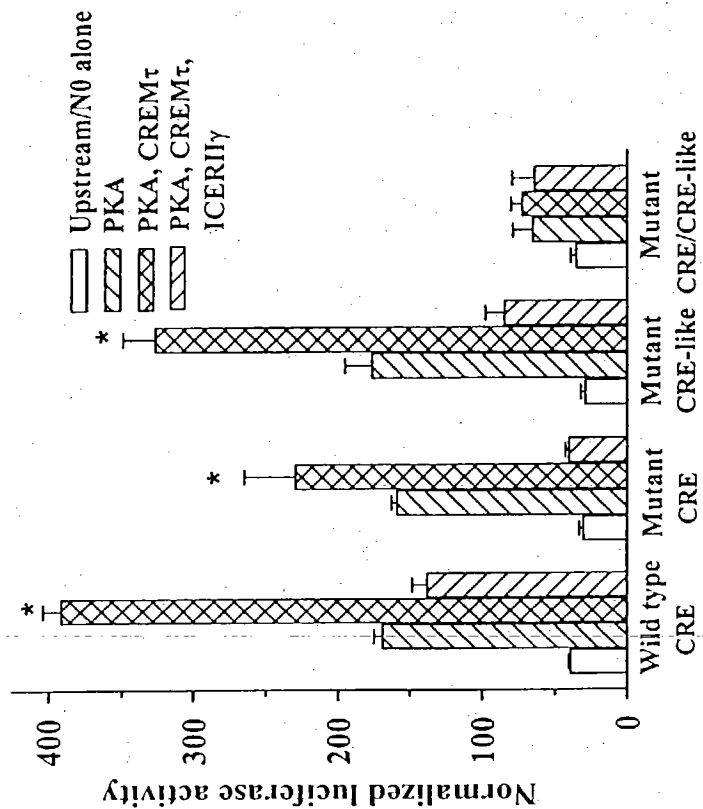


FIG. 5B

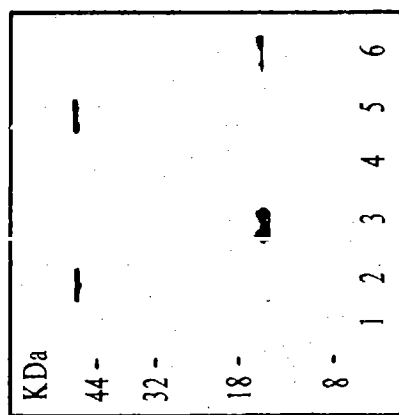


FIG. 5A

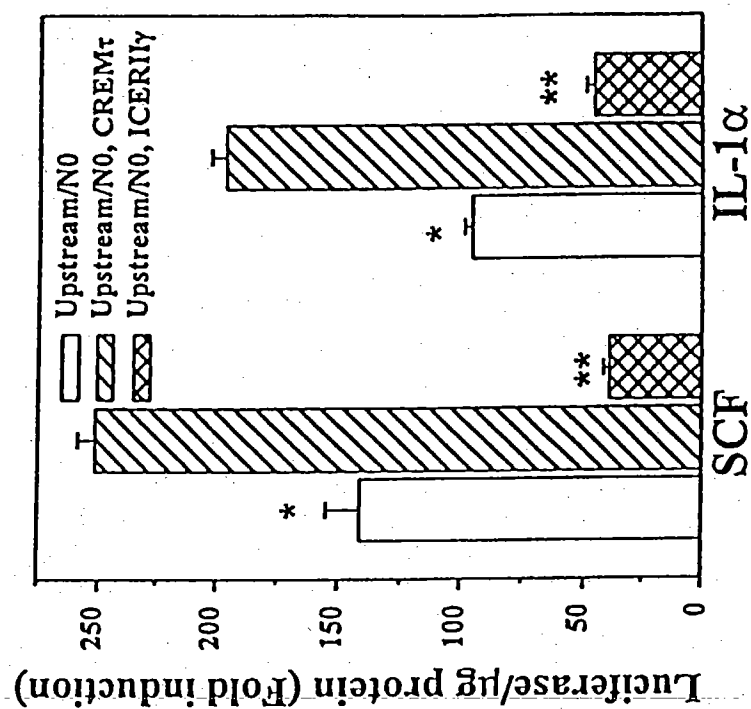


FIG. 6B

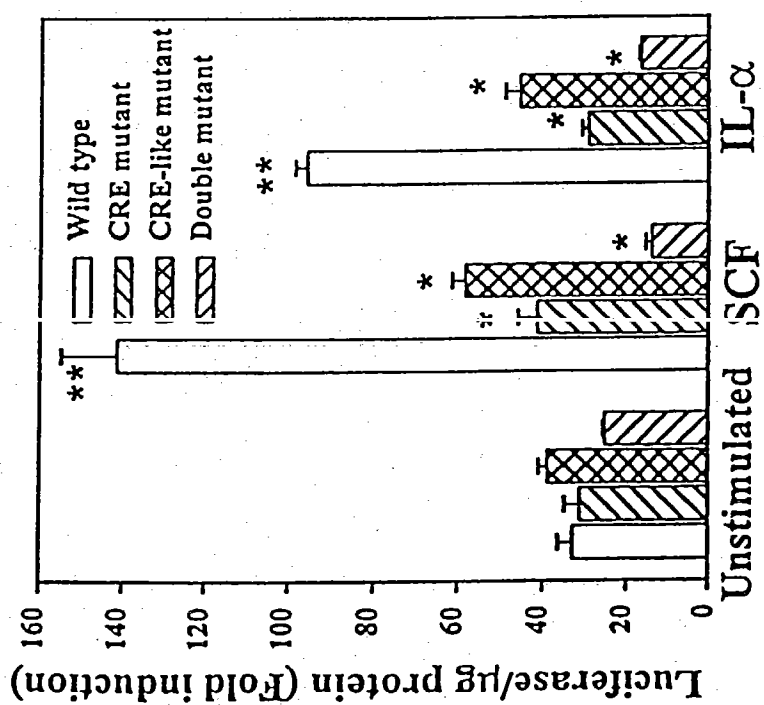
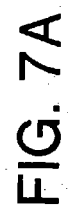
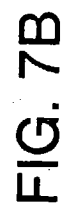


FIG. 6A



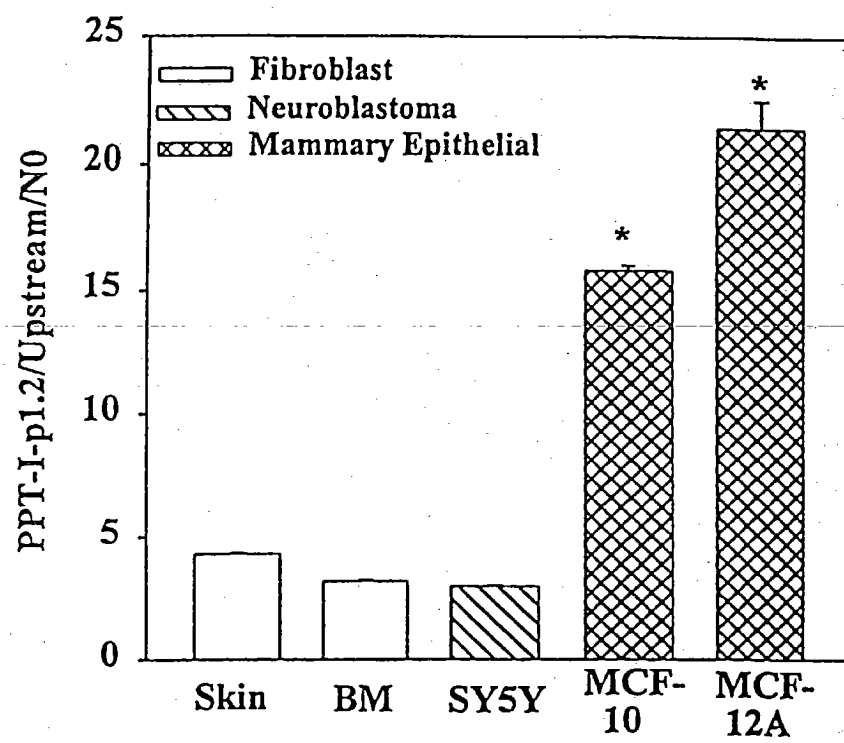


FIG. 8